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,
, TELECOMMUNICATION INFORMATION
, TELEPHONE: (650) 855-0555
, TELEFAX: (650) 845-4166
,
, INFORMATION FOR SEQ ID NO: 1495:
, SEQUENCE CHARACTERISTICS:
, LENGTH: 1642 base pairs
, TYPE: nucleic acid
, STRANDEDNESS: single
, TOPOLOGY: linear
, IMMEDIATE SOURCE:
, LIBRARY: GENBANK
, CLONE: G809486
, SEQUENCE DESCRIPTION: SEQ ID N
US-10-641-643-1485
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[illegible]

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RESULT 5
US-10-283-975A-302
; Sequence 302, Application US/10283975A
; Publication No. US20040110792A1
; GENERAL INFORMATION:
; APPLICANT: Ortho-Clinical Diagnostics, Inc.
; TITLE OF INVENTION: Methods For Assessing and Treating Leukemia
; FILE REFERENCE: CDS 293 PCT
; CURRENT APPLICATION NUMBER: US/10/283,975A
; CURRENT FILING DATE: 2002-10-30
; PRIOR APPLICATION NUMBER: 60/340,938
; PRIOR FILING DATE: 2001-10-30
; PRIOR APPLICATION NUMBER: 60/338,997
; PRIOR FILING DATE: 2001-10-30
; PRIOR APPLICATION NUMBER: 60/340,081
; PRIOR FILING DATE: 2001-10-30
; PRIOR APPLICATION NUMBER: 60/341,012
; PRIOR FILING DATE: 2001-10-30
; NUMBER OF SEQ ID NOS: 900
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 302
; LENGTH: 1642
; TYPE: DNA
; ORGANISM: HUMAN
US-10-283-975A-302

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Query Match	100.0%;	Score 348;	DB 18;	Length 1642;
Best Local Similarity	100.0%;	Pred. No. 2.3e-102;		
Matches 348;	Conservative	0;	Mismatches 0;	Indels 0;
Gaps 0;				

Qy	1	TT	C	A	G	A	G	C	G	G	G	C	G	G	G	C	G	C	C	T	G	G	G	A	A	G	A	G	A	C	T	G	T	G	T	G	C	A	G	A	T	T	60								
Db	- 373	TT	C	A	G	A	G	C	G	G	G	C	G	G	G	C	G	C	C	T	G	G	G	A	A	G	A	G	A	C	T	G	T	G	T	G	C	A	G	A	T	432									
Qy	61	A	A	C	G	T	C	A	T	A	T	G	T	G	T	A	A	T	G	T	G	G	G	A	A	G	A	T	T	G	G	T	G	T	G	T	C	A	G	T	C	A	A	A	G	T	C	120			
Db	433	A	A	C	G	T	C	A	T	A	T	G	T	G	T	A	A	T	G	T	G	G	G	A	A	G	A	T	T	G	G	T	G	T	G	T	C	A	G	T	C	A	A	A	G	T	C	492			
Qy	121	T	C	A	G	A	C	A	C	A	A	G	A	C	A	G	A	C	A	G	A	T	A	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	180				
Db	493	T	C	A	G	A	C	A	C	A	G	A	C	A	G	A	C	A	G	A	T	A	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	552					
Qy	181	C	G	G	A	G	T	C	A	G	A	A	T	C	T	G	A	A	G	A	A	C	A	C	A	G	A	A	C	G	A	A	C	A	G	T	G	G	C	C	C	C	C	C	C	C	240				
Db	553	C	G	G	A	G	T	C	A	G	A	A	T	C	T	G	A	A	G	A	A	C	A	C	A	G	A	A	C	G	A	A	C	A	G	T	G	G	C	C	C	C	C	C	C	C	612				
Qy	241	G	T	G	G	G	G	C	T	C	A	G	T	C	C	A	G	A	T	G	A	A	C	T	G	T	G	G	T	G	C	T	G	T	G	C	T	G	T	C	A	A	G	A	G	T	T	C	A	G	300
Db	613	G	T	G	G	G	G	C	T	C	A	G	T	C	C	A	G	A	T	G	A	A	C	T	G	T	G	G	T	G	C	T	G	T	G	C	T	G	T	C	A	A	G	A	G	T	T	C	A	G	672
Qy	301	C	A	G	C	C	C	G	T	G	A	C	C	T	C	C	A	A	C	A	G	A	G	T	G	G	G	C	A	T	G	T	C	C	C	C	G	A	T	G	T	C	A	348							
Db	673	C	A	G	C	C	C	G	T	G	A	C	C	T	C	C	A	A	C	A	G	A	G	T	G	G	G	C	A	T	G	T	C	C	C	G	A	T	G	T	C	A	720								

RESULT 6

US-10-825-282-13
; Sequence 13, Application US/10825282
; Publication No. US20040224389A1
; GENERAL INFORMATION:
; APPLICANT: 3921-1-1-1
; TITLE OF INVENTION: VIRAL VECTORS ENCODING APOPTOSIS-INDUCING PROTEINS AND
; METHOD OF INVENTION: METHODS FOR MAKING AND USING THE SAME
; FILE REFERENCE: 3921-1-1-1
; CURRENT APPLICATION NUMBER: 2004-04-14
; CURRENT FILING DATE: 2004-04-14
; PRIORITY APPLICATION NUMBER: US/09/456,357
; PRIORITY FILING DATE: 1999-12-08
; PRIORITY APPLICATION NUMBER: 60/134,416
; PRIORITY FILING DATE: 1999-05-17
; PRIORITY APPLICATION NUMBER: 09/087,195
; PRIORITY FILING DATE: 1998-05-29
; PRIORITY APPLICATION NUMBER: 08/378,507
; PRIORITY FILING DATE: 1995-01-26
; PRIORITY APPLICATION NUMBER: 08/250,478
; PRIORITY FILING DATE: 1994-05-27
; NUMBER OF SEQ ID NOS: 50
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
; LENGTH: 1642
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (130)..(756)
; FEATURES:
; 130-825-282-13

not a pure art-

Box #13 is same

no FADD - (p. 21 spec.
(MOR 1) (2910/825-282

	Query Match	100.0%	Score 348;	DB 18;	Length 1642;
	Best Local Similarity	100.0%;	Pred. No. 2.3e-102;		
	Matches 348;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0
QY	1	TTCAGCGGGGCGCGCGCGCGCGCTGTGGGAAGAAGACCTGTGTGCAGCATTT			60
Dδ	373	TTCAGCGGGGCGCGCGCGCGCGCTGTGGGAAGAAGACCTGTGTGCAGCATTT			432
QY	61	AACGTCTATGTGATAATGTGGGAAAGATTGGAGAAGCGTGGCTCGTCAGCTCAAAGTC			120
Dδ	433	AACGTCTATGTGATAATGTGGGAAAGATTGGAGAAGCGTGGCTCGTCAGCTCAAAGTC			492
QY	121	TCAGACACCAAGATCGACAGCATCGAGAGAGATACCCCGAACCTCACAGAGCGTG			180
Dδ	493	TCAGACACCAAGATCGACAGCATCGAGAGAGATACCCCGAACCTCACAGAGCGTG			552

QY 61 AACGTCATATGTGATAATGTGGGAAAGATTGGAGAGGCTGGCTCGTCAGCTCAAGTC 120
 Db 448 AACGTCATATGTGATAATGTGGGAAAGATTGGAGAGGCTGGCTCGTCAGCTCAAGTC 507
 QY 121 TCAGACACCAAGATCGACAGCATCGAGGACAGATACCCCGCAACCTCGACAGAGCGGTG 180
 Db 508 TCAGACACCAAGATCGACAGCATCGAGGACAGATACCCCGCAACCTCGACAGAGCGGTG 567
 QY 181 CGGGAGTCACTGAGATCTGAGAAACACAGAGAGGAGAACCGAACAGTGCGGCCACCTG 240
 Db 568 CGGGAGTCACTGAGATCTGAGAAACACAGAGAGGAGAACCGAACAGTGCGGCCACCTG 627
 QY 241 GTGGGGCTCTCAGGTCTCGCAGATGAACCTGGTGGCTGACCTGTACAGAGGTTTCAG 300
 Db 628 GTGGGGCTCTCAGGTCTCGCAGATGAACCTGGTGGCTGACCTGTACAGAGGTTTCAG 687
 QY 301 CAGGCCCGTGACCTCCAGAACAGGAGTGGGGCCATGTCCCGCATGTCA 348
 Db 688 CAGGCCCGTGACCTCCAGAACAGGAGTGGGGCCATGTCCCGCATGTCA 735

RESULT 11

US-10-723-860-5755
 ; Sequence 5755, Application US/10723860
 ; Publication No. US20040253606A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Aziz, Nacasha
 ; APPLICANT: Ginsburg, Wendy M.
 ; APPLICANT: Zlotnik, Albert
 ; TITLE OF INVENTION: Methods of Diagnosis of Soft Tissue Sarcoma, Compositions &
 ; FILE REFERENCE: 05982.0193.NPUS01
 ; CURRENT APPLICATION NUMBER: US/10/723,860
 ; CURRENT FILING DATE: 2003-11-26
 ; PRIOR APPLICATION NUMBER: 60/429,739
 ; PRIOR FILING DATE: 2002-11-26
 ; NUMBER OF SEQ ID NOS: 8393
 ; SOFTWARE: PatentIn version 3.2
 ; SEQ ID NO 5755
 ; LENGTH: 2288
 ; TYPE: DNA
 ; ORGANISM: Homo sapiens
 ; FEATURE:
 ; NAME/KEY: misc_feature
 ; LOCATION: (839)..(841)
 ; OTHER INFORMATION: n is a, c, g, or t
 ; FEATURE:
 ; NAME/KEY: misc_feature
 ; LOCATION: (843)..(846)
 ; OTHER INFORMATION: n is a, c, g, or t
 ; FEATURE:
 ; NAME/KEY: misc_feature
 ; LOCATION: (849)..(849)
 ; OTHER INFORMATION: n is a, c, g, or t
 ; FEATURE:
 ; NAME/KEY: misc_feature
 ; LOCATION: (852)..(859)
 ; OTHER INFORMATION: n is a, c, g, or t
 ; FEATURE:
 ; NAME/KEY: misc_feature
 ; LOCATION: (862)..(863)
 ; OTHER INFORMATION: n is a, c, g, or t
 ; FEATURE:
 ; NAME/KEY: misc_feature
 ; LOCATION: (865)..(866)
 ; OTHER INFORMATION: n is a, c, g, or t
 US-10-723-860-5755

Query Match 96.1%; Score 334.4; DB 18; Length 2288;
 Best Local Similarity 99.4%; Pred. No. 6.6e-98;
 Matches 346; Conservative 0; Mismatches 1; Indels 1; Gaps 1;

QY 1 TTCAGGCGGGCGCGCGCGCGCGCGCGCTGGGGAAGAACCTGTGTGCAGCATTT 60
 Db 334 TTCAGGCGGGCGCGCGCGCGCGCGCGCTGGGGAAGAACCTGTGTGCAGCATTT 392
 QY 61 AACGTCATATGTGATAATGTGGGAAAGATTGGAGAGGCTGGCTCGTCAGCTCAAGTC 120
 Db 393 AACGTCATATGTGATAATGTGGGAAAGATTGGAGAGGCTGGCTCGTCAGCTCAAGTC 452
 QY 121 TCAGACACCAAGATCGACAGCATCGAGGACAGATACCCCGCAACCTCGACAGAGCGGTG 180
 Db 453 TCAGACACCAAGATCGACAGCATCGAGGACAGATACCCCGCAACCTCGACAGAGCGGTG 512
 QY 181 CGGGAGTCACTGAGATCTGGAAGAACACAGAGAGGAGAACCGAACAGTGCGGCCACCTG 240
 Db 513 CGGGAGTCACTGAGATCTGGAAGAACACAGAGAGGAGAACCGAACAGTGCGGCCACCTG 572
 QY 241 GTGGGGCTCTCAGGTCTCGCAGATGAACCTGGTGGCTGACCTGTACAGAGGTTTCAG 300
 Db 573 GTGGGGCTCTCAGGTCTCGCAGATGAACCTGGTGGCTGACCTGTACAGAGGTTTCAG 632
 QY 301 CAGGCCCGTGACCTCCAGAACAGGAGTGGGGCCATGTCCCGCATGTCA 348
 Db 633 CAGGCCCGTGACCTCCAGAACAGGAGTGGGGCCATGTCCCGCATGTCA 680

RESULT 12

US-10-207-655-417
 ; Sequence 417, Application US/10207655
 ; Publication No. US20030118592A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Ledbetter, Jeffrey A.
 ; APPLICANT: Hayden-Ledbetter, Martha S.
 ; TITLE OF INVENTION: BINDING DOMAIN-IMMUNOGLOBULIN FUSION PROTEINS
 ; FILE REFERENCE: 390069.401C1
 ; CURRENT APPLICATION NUMBER: US/10/207,655
 ; CURRENT FILING DATE: 2002-07-25
 ; NUMBER OF SEQ ID NOS: 426
 ; SOFTWARE: PatentIn version 3.0
 ; SEQ ID NO 417
 ; LENGTH: 645
 ; TYPE: DNA
 ; ORGANISM: Artificial Sequence
 ; FEATURE:
 ; OTHER INFORMATION: fusion polynucleotide
 US-10-207-655-417

Query Match 48.6%; Score 169.2; DB 15; Length 645;
 Best Local Similarity 72.5%; Pred. No. 1.9e-44;
 Matches 219; Conservative 0; Mismatches 83; Indels 0; Gaps 0;

QY 1 TTCAGGCGGGCGCGCGCGCGCGCGCTGGGGAAGAACCTGTGTGCAGCATTT 60
 Db 259 TTCAGGCGGGCGCGCGCGCGCGCGCTGGGGAAGAACCTGTGTGCAGCATTT 318
 QY 61 AACGTCATATGTGATAATGTGGGAAAGATTGGAGAGGCTGGCTCGTCAGCTCAAGTC 120
 Db 319 GACATTTGTGTGACAAATGTGGGAGAGACTGGAAGAAGCTGCCCGGAGCTGAAGTG 378
 QY 121 TCAGACACCAAGATCGACAGCATCGAGGACAGATACCCCGCAACCTCGACAGAGCGGTG 180
 Db 379 TCTGAGGCCAAGATGATGGATTGAGGAGAGTACCCCGAAGTCTGAGTGAGCGGTA 438
 QY 181 CGGGAGTCACTGAGATCTGGAGAACACAGAGAGGAGAACCGAACAGTGCGGCCACCTG 240
 Db 439 AGGAGAGTCTGAAAGTCTGGAAGAATGCTGGAAGAAGAACGCTCGGTGGCCGAGTG 498
 QY 241 GTGGGGCTCTCAGGTCTCGCAGATGAACCTGGTGGCTGACCTGTGTACAGAGGTTTCAG 300
 Db 499 GTCAAGCGCTGCGGACCTGCAGGCTGAATCTGTGTGCTGACCTGTGTGGAGAGGCCAG 558
 QY 301 CA 302
 Db 559 GA 560

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RESULT 13
US-09-918-995-29039
; Sequence 29039, Application US/09918995
; Publication No. US20030073623A1
; GENERAL INFORMATION:
; APPLICANT: Hyseq, Inc.
; TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
; FILE REFERENCE: 20411-756
; CURRENT APPLICATION NUMBER: US/09/918,995
; PRIOR FILING DATE: 2001-07-30
; PRIOR APPLICATION NUMBER: US/09/235,076
; PRIOR FILING DATE: 1999-01-20
; NUMBER OF SEQ ID NOS: 38054
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 29039
; LENGTH: 474
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(474)
; OTHER INFORMATION: n = A,T,C or G
US-09-918-995-29039

Query Match      27.6%; Score 96.2; DB 10; Length 474;
Best Local Similarity 92.7%; Pred. No. 8.3e-21;
Matches 101; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

QY      240 GGTGGGGCTCTCAGGTCTCCAGATGAACCTGGTGGCTGACCTGGTCAAGAGGTCA 299
Db      54 GGAGGAATTCCTCAGGTCTCCAGATGAACCTGGTGGCTGACCTGGTCAATAGGTCA 113

QY      300 GCAGGGCCGTGACCTCCAGAACAGGAGTGGGGCCATGTCCCGATGTCA 348
Db      114 GCAGGCCCGTGACCTCCAGAACAGGAGTGGGGCCATGTCCCGATGTCA 162

RESULT 14
US-09-864-761-2348/c
; Sequence 2348, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharron G.
; APPLICANT: Rank, David R.
; APPLICANT: Hanzel, David K.
; APPLICANT: Chen, Wensheng
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
; FILE REFERENCE: Aeomica-X-1
; CURRENT APPLICATION NUMBER: US/09/864,761
; CURRENT FILING DATE: 2001-05-23
; PRIOR APPLICATION NUMBER: US 60/180,312
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: US 60/207,456
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: US 09/632,366
; PRIOR FILING DATE: 2000-08-03
; PRIOR APPLICATION NUMBER: GB 24263.6
; PRIOR FILING DATE: 2000-10-04
; PRIOR APPLICATION NUMBER: US 60/236,359
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: PCT/US01/00666
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00667
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00664
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00669
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00665
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; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00668
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00663
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00662
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00661
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00670
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: US 60/234,687
; PRIOR FILING DATE: 2000-09-21
; PRIOR APPLICATION NUMBER: US 09/608,408
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: US 09/774,203
; PRIOR FILING DATE: 2001-01-29
; NUMBER OF SEQ ID NOS: 49117
; SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 2348
; LENGTH: 485
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: MAP TO AL035661.16
; OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 22
; OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 7.2
; OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 3.6
; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 8.3
; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 3.4
; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 5.8
; OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 3.4
; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 18
; OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 4.8
; OTHER INFORMATION: EXPRESSED IN HBL100, SIGNAL = 3.1
US-09-864-761-2348

Query Match      11.8%; Score 41; DB 9; Length 485;
Best Local Similarity 53.4%; Pred. No. 0.0069;
Matches 86; Conservative 0; Mismatches 75; Indels 0; Gaps 0;

QY      155 ACCCCCGCAACCTGACAGAGCGTGTCCGGAGTCACTGAGAACTCGGAGAACACACAGAGA 214
Db      452 ACCAAGGACTCCGGGGTGGCGCGTGTGAGAGCTGAAGAAATAGTGGATGAGCGGTGTA 393

QY      215 AGGAGAACGCAACAGTGGCCACCTGGTGGGGCTCTCAGGTCTCCAGATGAACCTGG 274
Db      392 AGCACTGCCCCACCGTGCAGCATGTCTGTGGTCTCACAGGACAGACACAAAGTCCACA 333

QY      275 TGGCTGACCTGGTACAAAGAGTTTCAGAGCGCCCGTGCACCTC 315
Db      332 TGGGGGATCTGGACGCTCCCGCTGAGCAGCGTGGGTACCTC 292

RESULT 15
US-10-024-298A-163
; Sequence 163, Application US/10024298A
; Publication No. US20030143540A1
; GENERAL INFORMATION:
; APPLICANT: ASAH KASEI KABUSHIKI KAISHA
; APPLICANT: AKIO MATSUDA
; APPLICANT: Goichi HONDA
; APPLICANT: Shuji MURAMATSU
; APPLICANT: Yukiko NAGANO
; TITLE OF INVENTION: NP-K B Activating Gene
; FILE REFERENCE: 1254-0191P
; CURRENT APPLICATION NUMBER: US/10/024,298A
; CURRENT FILING DATE: 2003-04-08
; PRIOR APPLICATION NUMBER: 60/314,385
; PRIOR FILING DATE: 2001-08-24
; PRIOR APPLICATION NUMBER: 60/278,641
; PRIOR FILING DATE: 2001-03-26
; PRIOR APPLICATION NUMBER: 60/258,315
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; PRIOR FILING DATE: 2000-12-28
; PRIOR APPLICATION NUMBER: JP254018/2001
; PRIOR FILING DATE: 2001-08-24
; PRIOR APPLICATION NUMBER: JP0088912/2001
; PRIOR FILING DATE: 2001-03-26
; PRIOR APPLICATION NUMBER: JP402288/2000
; PRIOR FILING DATE: 2000-12-28
; NUMBER OF SEQ ID NOS: 182
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 163
; LENGTH: 4031
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (91)...(2649)
US-10-024-298A-163
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Query Match      11.4%; Score 39.8; DB 15; Length 4031;
Best Local Similarity 59.2%; Pred. No. 0.029;
Matches 87; Conservative 0; Mismatches 57; Indels 3; Gaps 1;

QY      129 CAAGATCGACAGCATCGAGGACAGATACCCCGCAACCTGACAGAGCGTGTGCGGAGTGC 188
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db      234 CAACATCTACACCTTCAACCCACACCCGTGACCCCGCAACCGGACCGAGGGTGTGCGAGTGC 293
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY      189 ACTGAGAATCTGGAAGAAC---ACAGAGAAGGAGAACGCAACAGTGGCCACCTGTGTGGG 245
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db      294 TGTGATGTCCTGAAACAAGCAGAAAGGGCGCCCTTTGCTGTGTCGCCAGAAAGGA 353
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY      246 GGCCTCTCAGGTCCTGCCAGATGAACCT 272
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db      354 GGCCTGTGTGTCCTTCCAGGTGCCCT 380
      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
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Search completed: February 11, 2005, 20:40:03
Job time : 390 secs